

PSRM_Replication.R

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```
#PSRM Replication Files
#
#The Population Ecology of Interest Groups and Counter-Mobilization:
#Reproductive Rights Organizations in the United States, 1920-1985.
#
#Tristan M. Hightower (2023)

#Change working directory for data loading

setwd("~/Desktop/PSRM Replication")

#Load libraries
library(readr)
library(imputeTS)

## Registered S3 method overwritten by 'quantmod':
##   method           from
##   as.zoo.data.frame zoo

library(readxl)
library(MASS)
library(stargazer)

##
## Please cite as:
## Hlavac, Marek (2022). stargazer: Well-Formatted Regression and Summary Statistics Tables.
## R package version 5.2.3. https://CRAN.R-project.org/package=stargazer

library(tidyverse)

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.1    v purrr      1.0.1
## v forcats   1.0.0    v stringr   1.5.0
## v ggplot2   3.4.3    v tibble    3.2.1
## v lubridate 1.9.2    v tidyr     1.3.0

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
## x dplyr::select() masks MASS::select()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors

library(growthcurver)
```

```

#Load Data
groups <- read_csv("PSRM_Replication.csv")

## Rows: 66 Columns: 21
## -- Column specification -----
## Delimiter: ","
## dbl (21): Year, Choice_founding, Choice_density, Choice_densitysq, Life_foun...
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

#Descriptive Figure 1

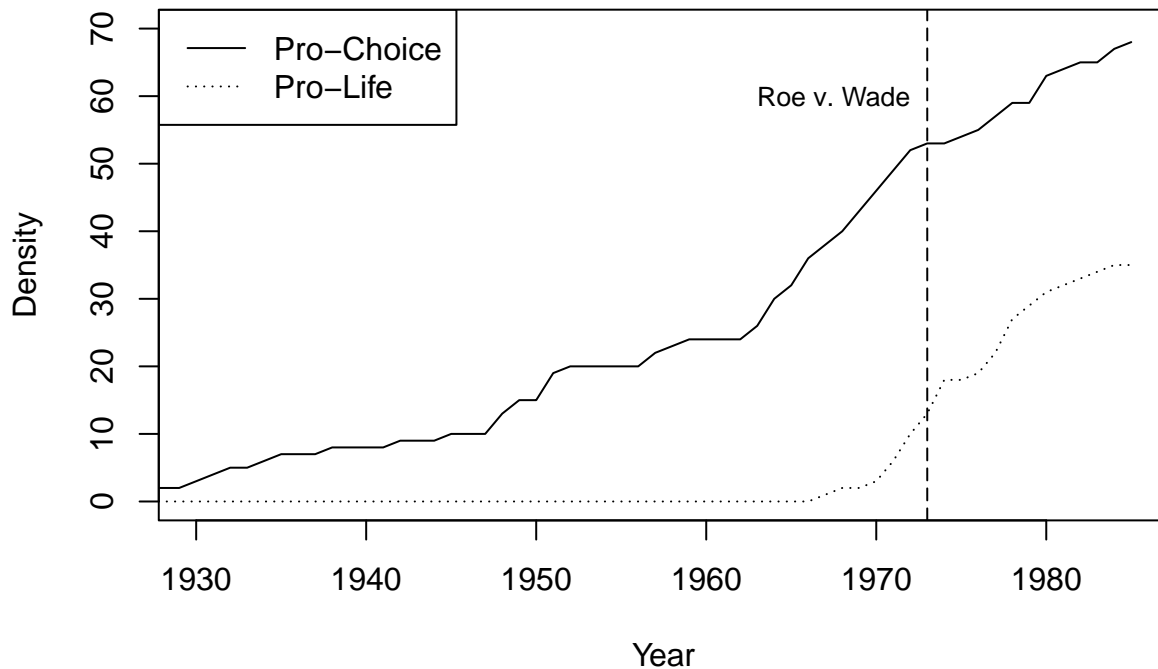
plot(groups$Year,groups$Choice_density, type = "l", xlab = "Year",
      ylab = "Density", xlim = c(1930,1985), ylim = c(0,70))

lines(groups$Year, groups$Life_density, lty = 3)

legend("topleft",
      legend=c("Pro-Choice", "Pro-Life"),
      lty=c(1,3))

abline(v=1973, lty = 5)
text(1967.5,60,"Roe v. Wade", cex = .8)

```



```

#Run Models
m1<-glm.nb(groups$Life_founding~groups$Life_density+groups$Life_densitysq+
           groups$Choice_density+groups$Choice_densitysq+groups$Abortion_ratio+
           groups$M_BA+groups$W_BA)

## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached

```

```
## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached
m2<-glm.nb(groups$Choice_founding~groups$Choice_density+groups$Choice_densitysq+
           groups$Life_density+groups$Life_densitysq+groups$Abortion_ratio+groups$M_BA+groups$W_BA)
```

```
## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached
```

```
## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached
```

```
#Create Table 1

stargazer(m1,m2, type = "text",
          column.labels = c("Anti-Abortion","Abortion-Rights"),
          dep.var.labels.include = F, dep.var.caption = "DV: Group Formation",
          covariate.labels = c("Anti-Abortion", "Anti-Abortion Sq.",
                               "Abortion Rights", "Abortion Rights Sq.",
                               "Abortion Ratio", "Male BA Degrees","Female BA Degrees"),
          notes = "Negative binomial regression with standard errors in parentheses.")
```

```
##
## =====
##                               DV: Group Formation
##                               -----
##                               Anti-Abortion           Abortion-Rights
##                               (1)                   (2)
##                               -----
## Anti-Abortion                0.113                 -0.609**
##                               (0.258)              (0.299)
##
## Anti-Abortion Sq.            0.003                 0.013***
##                               (0.006)              (0.005)
##
## Abortion Rights              1.181*                0.283**
##                               (0.640)              (0.118)
##
## Abortion Rights Sq.         -0.011*                -0.001
##                               (0.006)              (0.001)
##
## Abortion Ratio               0.001                 0.018*
##                               (0.012)              (0.011)
##
## Male BA Degrees             -1.122                -0.520**
##                               (1.296)              (0.238)
##
## Female BA Degrees            0.293                 -0.387
##                               (1.256)              (0.391)
##
## Constant                    -18.473               1.611
##                               (15.318)              (1.236)
##
## -----
## Observations                 66                   66
```

```
## Log Likelihood                -27.960                -74.953
## theta                        25,229.360 (528,093.700)    13,587.290 (212,777.800)
## Akaike Inf. Crit.            71.920                165.907
## =====
## Note:                        *p<0.1; **p<0.05; ***p<0.01
##                               Negative binomial regression with standard errors in parentheses.
```

#Predictions

```
predictions <- data.frame("Choice_density" = 0:65, "Life_density" = 0:65, "Abortion_ratio" = mean(groups))
predictions$Life_densitysq <- (predictions$Life_density)^2
predictions['Choice_foundings'] <- NA
predictions$Choice_densitysq <- (predictions$Choice_density)^2

predictions['Life_foundings'] <- NA

predictions$Choice_foundings <- predict(m2, newdata = predictions)
predictions$Life_foundings <- predict(m1, newdata = predictions)
```

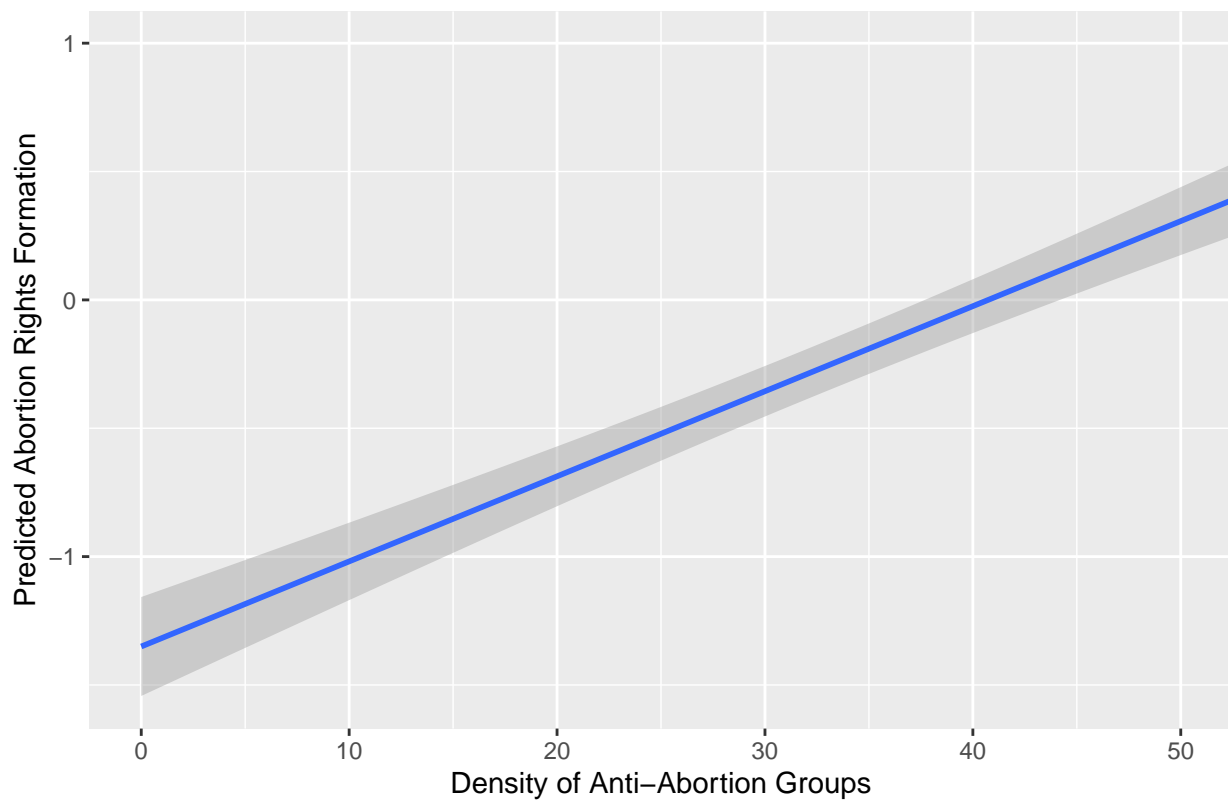
#Plotting Predictions

#Figure 2, publication

```
ggplot(predictions, aes(Life_density, Choice_foundings)) +
  geom_smooth(level = .9, method = "lm") +
  xlab("Density of Anti-Abortion Groups") +
  ylab("Predicted Abortion Rights Formation") +
  labs(title = "Pred. Abortion-Rights Foundings Relative to Anti-Abortion Density") +
  coord_cartesian(xlim = c(0, 50))
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

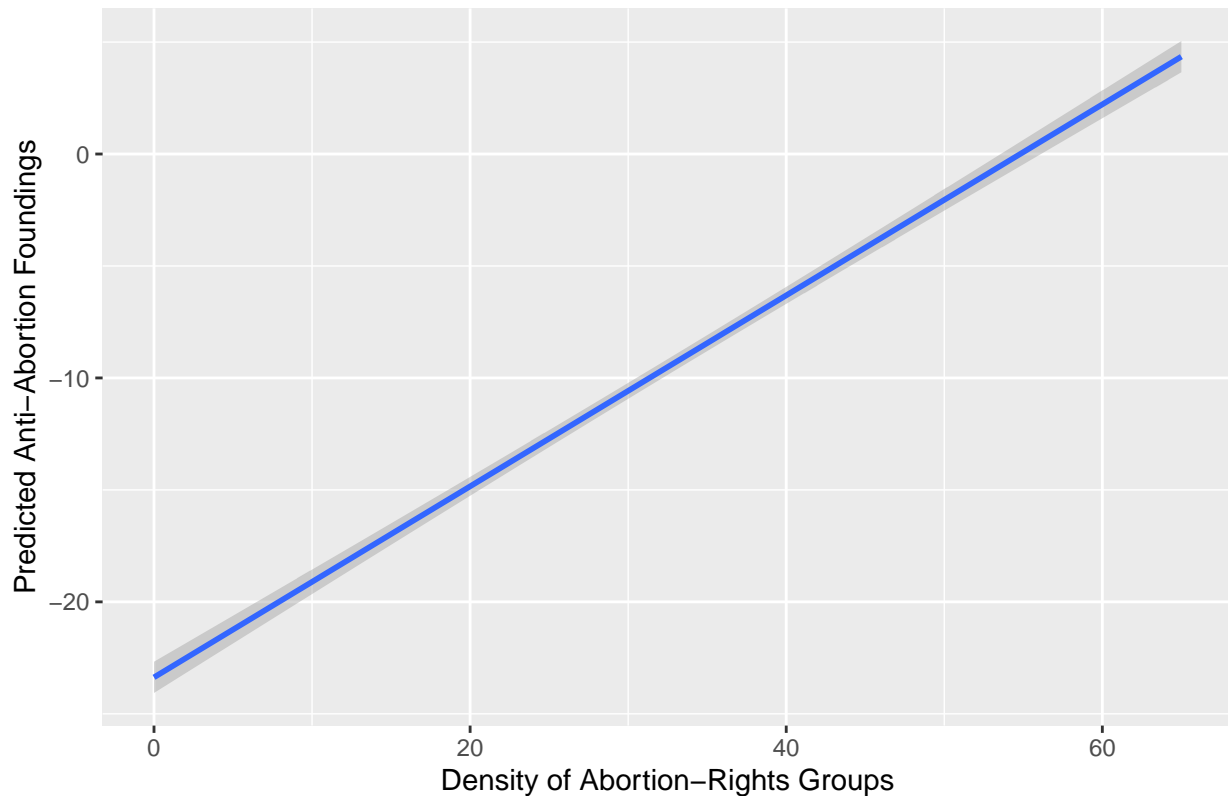
Pred. Abortion-Rights Foundings Relative to Anti-Abortion Density



```
#Figure 3, publication  
ggplot(predictions, aes(Life_density, Life_foundings)) +  
  geom_smooth(level = .9, method = "lm") +  
  xlab("Density of Abortion-Rights Groups")+  
  ylab("Predicted Anti-Abortion Foundings")+  
  labs(title = "Pred. Anti-Abortion Foundings Relative to Abortion-Rights Density")
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

Pred. Anti-Abortion Foundings Relative to Abortion-Rights Density



```
###Appendices###
```

```
#Creating figures 4 and 5 using growthcurver 'Summarize_Growth' function
```

```
#creating blank dataset into 'd2'
```

```
groups$time<-c(1:66)
```

```
d2<-groups$time
```

```
d2<-as.data.frame(d2)
```

```
colnames(d2)<-"time"
```

```
d2$choice<-groups$Choice_density
```

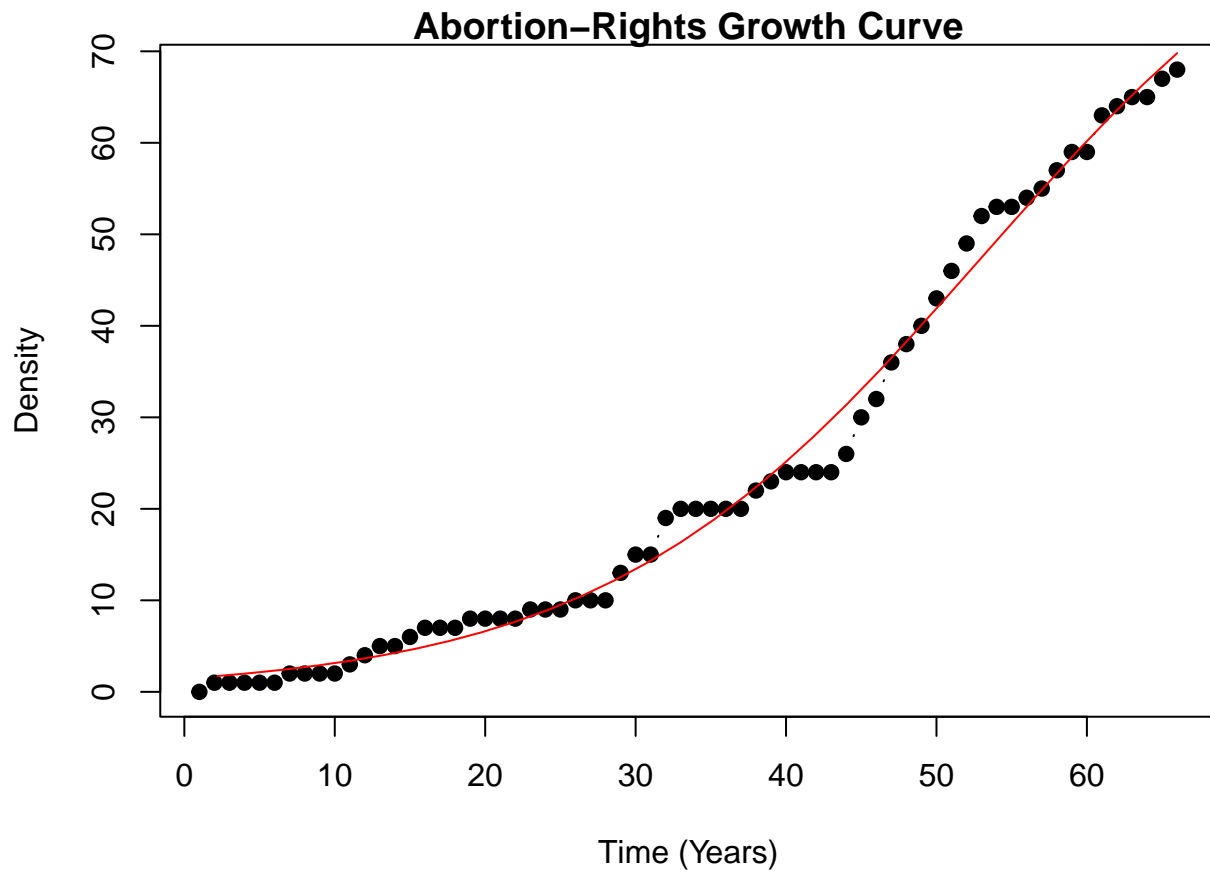
```
d2$life<-groups$Life_density
```

```
#running model for abortion rights
```

```
gc_fit_choice<-SummarizeGrowth(d2$time,d2$choice)
```

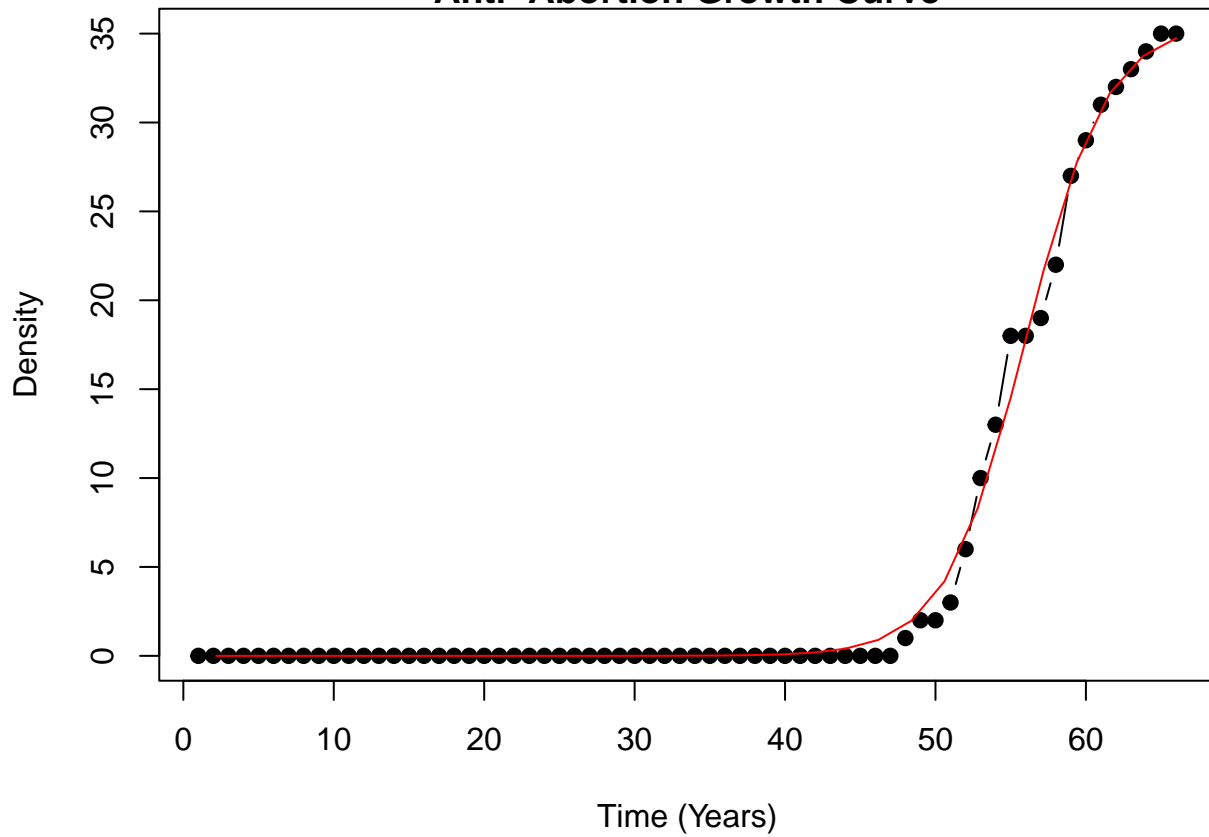
```
#Creating Figure 4
```

```
plot(gc_fit_choice, xlab = "Time (Years)", ylab = "Density", main = "Abortion-Rights Growth Curve")
```



```
#running model for anti-abortion  
gc_fit_life<-SummarizeGrowth(d2$time,d2$life)  
  
#Creating Figure 5  
plot(gc_fit_life, ylab="Density", xlab = "Time (Years)", main = "Anti-Abortion Growth Curve")
```

Anti-Abortion Growth Curve



```

#Creating Table 2
groups2<-groups[groups$Year>1966,]
m3<-glm.nb(groups2$Life_founding~groups2$Life_density+groups2$Life_densitysq+groups2$Abortion_ratio)

## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached

## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached
m4<-glm.nb(groups2$Life_founding~groups2$Choice_density+groups2$Choice_densitysq+groups2$Abortion_ratio)

## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached

## Warning in theta.ml(Y, mu, sum(w), w, limit = control$maxit, trace =
## control$trace > : iteration limit reached
stargazer(m3,m4,dep.var.caption = "DV: Anti-Abortion Group Formation",dep.var.labels.include = F,
  type = "text", covariate.labels = c("Anti-Abortion", "Anti-Abortion Sq.", "Abortion Rights",
  "Abortion Rights Sq.", "Abortion Ratio"),
  omit = c("Constant"), omit.stat = "theta", notes = "Negative binomial regression with SEs in p

##
## =====
##                   DV: Anti-Abortion Group Formation
##                   -----
##                   (1)                   (2)
##

```

```

## -----
## Anti-Abortion          0.356
##                       (0.231)
##
## Anti-Abortion Sq.     -0.007**
##                       (0.004)
##
## Abortion Rights       0.844**
##                       (0.420)
##
## Abortion Rights Sq.   -0.008**
##                       (0.003)
##
## Abortion Ratio        -0.007          -0.00001
##                       (0.008)          (0.004)
## -----
## Observations          19              19
## Log Likelihood        -29.701         -29.526
## Akaike Inf. Crit.     67.403          67.051
## =====
## Note:                  *p<0.1; **p<0.05; ***p<0.01
##                       Negative binomial regression with SEs in parentheses.

```